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## DESCRIPTION

RECOMBINANT HERPESVIRUS AND USES THEREOF

## 5 Technical Field

The present invention relates to a recombinant herpesvirus having DNA that encodes the gB gene of infectious laryngotracheitis virus (hereinafter referred to as ILTV), and uses thereof. More specifically, it  
10 relates to a recombinant herpesvirus having a partial sequence of the gB gene of ILTV that can be stably present in the recombinant, and a vaccine against infectious laryngotracheitis virus.

## 15 Background Art

Infectious laryngotracheitis is caused by the infection of ILTV. ILTV infects avians such as chickens, pheasants, peacocks and turkeys. Characteristic features of onset in chickens include the appearance of  
20 respiratory symptoms, elevation of body temperature and anorexia and the like, serious coughing, expectoration of sputums. When egg-laying chickens are infected, the rate of laying eggs starts to decrease about four days after the onset of the disease, and takes about one month  
25 before normal egg-laying returns. Furthermore, increases in death rate due to mixed infection of ILTV and other pathogens have been reported, and infectious laryngotracheitis inflicts huge economic losses on the poultry industry.

30 For the prevention of infectious laryngotracheitis, dried live vaccines or frozen live vaccines from attenuated vaccine strains have conventionally been used. However, the effect of immunization varies with the breeding environment, the breeding density, the  
35 inoculation method and the like. Furthermore, there are also risks that vaccination may cause slight symptoms in the respiratory tract and faulty methods or amounts of

inoculation may lead to the onset of the disease. In some areas, there are reports of diseases caused by the reverted pathogenicity of vaccine strains, and thus the development of safe and effective vaccines is being sought.

In order to overcome the problems, recently, vaccines comprising as an active ingredient a recombinant virus vector have been developed by the recombinant technology. With regard to ILTV, the use of fowlpox virus (hereinafter referred to as FPV) as the virus vector has been investigated and it is commercially available in the USA (BIOMUNE, trade name VECTORMUNE FP-LT(+AE)).

ILTV is a causative virus of infectious laryngotracheitis. ILTV is one of the herpesviruses and the virus genome consists of a double stranded DNA comprising about 160,000 base pairs. There are known the thymidine kinase gene (Griffin et al., J. Gen. Virol. 71:841, 1990), gp60 gene (Kongsuwan et al., Virus Genes 7:297-303, 1993), the capsid p40 gene (Griffin et al., Nucleic Acids Res. 18:366, 1990), the glycoprotein B (gB) gene (Poulsen et al., Virus Genes 5:335-347, 1991; Griffin et al., J. Gen. Virol. 72:393-398, 1991; U.S. Pat. No. 5,443,831), the glycoprotein C (gC) gene (Kingsley et al., Virology 203:336-343, 1994), the RR2 gene (Griffin et al., J. of General Virol. 70:3085-3089, 1989), the UL32 gene (International Patent Publication WO98/07866), and the like.

The open reading frame of gB gene of ILTV has a full-length of 2613 bp (873 amino acids), and it is reported in U.S. Pat. No. 5,443,831 etc. that a recombinant FPV having the full-length gene integrated therein exhibits an effect as a vaccine.

Poxviruses such as FPV rapidly grow in the host and after expressing the antigen protein, they are completely expelled from the host's immune system. However, as immunity is memorized and boosted after expelling the

virus, the poxvirus is suitable for use as the host for vaccine. On the other hand, herpesviruses such as turkey herpesvirus (hereinafter referred to as HVT) do not rapidly grow in the host, and latent infection lasts for a long time. During this latency, herpesviruses continue to stimulate the immune system of the host. The use of recombinant HVT as a virus vector using such a characteristic is being investigated.

Although PCT Japanese National Publication No. 4-501658 (EP 434721) described that a recombinant HVT comprising HVT, into which an ILTV antigen gene has been inserted, has been constructed in fact the recombinant HVT has not been constructed.

In addition, in Japanese Unexamined Patent Publication No. 2001-000188, a recombinant HVT, into which two genes, i.e., a full-length gB gene of ILTV and an UL32 gene have been inserted, has been constructed. It has been confirmed that this recombinant HVT expresses in vitro a protein corresponding to the inserted gene using a immunofluorescent assay method, but an effect as a vaccine has not been confirmed.

#### Disclosure of the Invention

Under the circumstances of the conventional technology, the present inventors have attempted to purify, by subculturing, a recombinant herpesvirus wherein a DNA molecule, in which a promoter was ligated upstream to the full-length gB gene of ILTV, has been inserted into the herpesvirus genome. However, the purification was impossible because of the deletion of ILTV gB gene. When the deletion of the gB gene occurs, the recombinant herpesvirus does not function as an anti-ILTV vaccine. Furthermore, ILTV and HVT are both herpesviruses, and HVT per se has the gB gene as an essential gene.

Thus, as gB genes may compete with each other at the time of forming virus particles, the present inventors

thought that it would be important to avoid completion and to retain antigenicity by deleting the membrane anchor portion and the cytoplasmic domain of the gB gene product of ILTV so as to change the gB protein of ILTV from a membrane protein to a secretary protein.

Based on this idea, after intensive and extensive study in order to obtain recombinant herpesviruses that are stable during subculturing, the present inventors have found that a vaccine effect can be obtained by shortening the gB gene to a predetermined length in addition to deleting the membrane anchor portion and the cytoplasmic domain, and a higher vaccine effect can be obtained by ligating the shortened gB gene to a specific additive sequence, and therefore have completed the present invention.

Thus, in accordance with the present invention, there is provided a recombinant herpesvirus (excluding infectious laryngotracheitis virus) having a DNA that encodes a polypeptide comprising 429 amino acids at the amino terminal end of a protein encoded by the gB gene of infectious laryngotracheitis virus or a polypeptide in which one or a plurality of amino acids have been deleted, added, or substituted in said polypeptide. Furthermore, there is provided a vaccine against infectious laryngotracheitis virus comprising as an active ingredient said recombinant herpesvirus.

#### Brief Explanation of the Drawings

Fig. 1 is a drawing that compares FW050 with the original homology vector.

Fig. 2 is a schematic diagram of the homology vector.

#### Best Mode for Carrying out the Invention

The present invention will now be explained in detail hereinbelow.

#### DNA

DNA for use in the present invention is one (hereinafter referred to as the partial gB gene) that encodes a partial peptide comprising 429 amino acids at the amino terminal end of a protein (hereinafter referred to as the gB protein) encoded by the gB gene of ILTV. The amino acid sequence of this partial peptide may include the deletion, addition or substitution of one or a plurality of amino acids.

As a specific example of DNA for use in the present invention, there can be mentioned a DNA that encodes a sequence of 429 amino acids set forth in SEQ ID NO: 2, and as a specific example, there can be mentioned a DNA of a nucleotide sequence set forth in SEQ ID NO: 3.

As ILTV that can be an origin of the gB gene, there can be mentioned, for example, the NS-175 strain (the strain number VA0204 of the Catalog of Cultures for Animal Hygiene, the Japanese Association of Veterinary Biologics), the CE strain (Kouda, Azabu Veterinary College Research Report, 31, 133-202, 1976), the SA-2 strain (Johnson et al., Arch. Virol., 119, 181-198, 1991), the highly toxic field isolate 632 strain (Keeler et al., Avian Diseases, 35, 920-929, 1991), and the USDA challenge strain (Poulsen et al., J. General. Virol., 78, 2945-2951, 1997).

The gB gene is not specifically limited, and any gene encoding the gB protein derived from ILTV may be used, and there can be mentioned, for example, the gB gene (GeneBank ACC. No. X56093) derived from the highly toxic field isolate 632 strain and the gB gene (GeneBank ACC. No. M64927) derived from the SA2 strain.

The vaccine effect can be enhanced by ligating a DNA (hereinafter referred to as the additive DNA) that encodes the amino acid sequence set forth in SEQ ID NO: 4 in frame with the 3'-end of the above-mentioned gB gene. One or a plurality of amino acids of the amino acid sequence set forth in SEQ ID NO: 4 may be deleted, added or substituted.

The linking of the additive DNA in a frame with the partial gB gene is both important and preferred in order to enhance the vaccine effect.

5 The amino acid sequence set forth in SEQ ID NO: 4  
neither has a sequence that has some function nor has a  
sequence that is estimated to have some function. Among  
the nucleic acid sequence 120 bp (not containing the stop  
codon) (SEQ ID NO: 5) encoding the amino acid sequence  
set forth in SEQ ID NO: 4, 87 bp is derived from the SV40  
10 polyadenylation signal, 16 bp from the UL45 of HVT, 13 bp  
from the artificially introduced SfiI sequence, and 4 bp  
is from the flanking sequence.

The method of ligating the partial gB gene and the  
additive DNA in a frame is not specifically limited, and  
15 a general method described in a textbook (Sambrook, J.,  
Russel D.W., ed., Molecular Cloning, Third Ed., Cold  
Spring Harbor Laboratory Press etc.) on gene recombinant  
technology can be used. As the general method, there can  
be mentioned a method of ligating with restriction enzyme  
20 cleavage sites. When restriction enzyme recognition  
sites are not available, such sites may be introduced by  
polymerase chain reaction (PCR), in vitro mutation or the  
like.

#### Recombinant herpesvirus

25 The recombinant herpesvirus of the present invention  
must employ a herpesvirus other than ILTV as the parent  
virus. The parent herpesvirus may be any herpesvirus  
that infects mammals or avians. However, as the  
integrated gene cannot stably exist in ILTV, ILTV cannot  
30 be used as the parent virus. When a vaccine for avians  
is to be obtained, it is preferred to select Marek's  
disease virus as the parent herpesvirus. There are three  
types of Marek's disease virus: type 1, 2 and 3, and any  
type can be selected in accordance with the present  
35 invention. A Marek's disease virus may be naturally  
obtained, or is available from ATCC etc., with or without  
cost, and non-pathogenic ones are preferred. As such

viruses, there can be mentioned, for example, the CV1988 (Rispens) strain as Marek's disease virus type 1, the SB-1 strain as the Marek's disease virus type 2, and FC126 (ATCC VR-584B), PB-THV1, H-2, YT-7 and HPRS-26 as Marek's disease viruses type 3.

The method of constructing a recombinant herpesvirus is not specifically limited, and a recombinant plasmid having an additive DNA ligated as desired to the above-mentioned partial gB gene may be used, which may be inserted into a parent herpesvirus by homologous recombination. At this time, the partial gB gene is located at a position that is under the control of a foreign or endogenous promoter.

As a plasmid for use as a recombinant plasmid (hereinafter referred to as homology vector), any commonly used one can be used and includes, for example, pBR322, pBR325, pBR327, pBR328, pUC8, pUC18 and pUC19.

The construction of a homology vector may be performed according to a standard method using restriction enzyme sites owned by the vector.

The homology vector is a plasmid in which, usually, the promoter and the polyA signal are added to the above-mentioned partial gB gene which is then inserted into a region nonessential for the growth of the recombinant herpesvirus.

The method of constructing the homology vector of the present invention will be explained hereinbelow.

When DNA is to be integrated into a recombinant herpesvirus, it is generally integrated so that the DNA molecule of the present invention is located under the control of a regulatory gene (promoter) in order to obtain high expression. The promoter may be any commonly used one that operates in an eukaryotic cell, and may be derived from an eukaryotic cell or from a virus. Specific examples of the promoter include the thymidine kinase promoter of herpesvirus (Ross et al., J. Gen. Virol. 74:371-377, 1993), the gB protein promoter of

turkey herpesvirus (HVT) and Marek's disease virus (MDV) type 1 (Ross et al., J. Gen. Virol. 74:371-377, 1993), the IE promoter of human cytomegalovirus (HCMV) (Stinski et al., J. Virol. 55:431-441, 1985), the SV40 promoter  
5 (Gunning et al., Proc. Natl. Acad. Sci. U.S.A. 84:4831-4835, 1987), the human  $\beta$ -acting promoter (Gunning et al., Proc. Natl. Acad. Sci. U.S.A. 84:4831-4835, 1987), the chicken  $\beta$ -actin promoter (Kost et al., Nucleic Acids Res. 11:8287-8301, 1983), the LTR promoter of Rous sarcoma  
10 virus (RSV) (Greuel et al., Virology 177:33-43, 1990), the Pec promoter (Japanese Unexamined Patent Publication (Kokai) No. 2001-000188) and the like.

By adding an enhancer which is a transcription-activating factor in addition to the promoter, a further  
15 effective expression may be expected (Stinski et al., J. Virol. 55:431-441, 1985). As the enhancer, part of the cytomegalovirus-derived promoter may be illustrated and, generally, the positional relationship to the inserted gene is not limited. As the promoter of this type, there  
20 can be illustrated the Pec promoter described in Japanese Unexamined Patent Publication (Kokai) No. 2001-000188.

Furthermore, by adding a polyadenylation signal downstream to the additive DNA, a specifically high expression may be expected.

25 As the polyadenylation signal, there can be illustrated a polyA signal such as SV40 (Gunning et al., Proc. Natl. Acad. Sci. U.S.A. 84:4831-4835, 1987) and the polyA signal of UL46h, UL47h and UL49h of Marek's disease virus (MDV) type 1 (Yanagida et al., J. Gen. Virol.  
30 74:1837-1845, 1993).

As the gene regions nonessential for the growth of herpesvirus in the case of Marek's disease virus (MDV) type 1, type 2 and type 3 (type 3 is the turkey herpesvirus), there can be illustrated the TK region  
35 (Ross et al., the 16th International Herpes Workshop, 1991), the US10 region (Sakaguchi et al., Vaccine 12:953-957, 1994), the US2 region (Sondermeijer, P.J. et al.,



Vaccine 11:349-358, 1993), the region between UL44 and 45 and between UL45 and 46 described in Japanese Unexamined Patent Publication (Kokai) No. 11-192093, and the like.

5 The homology vector may be constructed by inserting a foreign gene such as the partial gB gene or the additive DNA, an exemplary DNA of the present invention. The length of the nonessential region into which foreign genes such as DNA of the present invention is inserted is not specifically limited, but 10 bp or more forward or  
10 behind the foreign gene insertion site, preferably 100 bp or more, and more preferably 500 bp or more of bases derived from the nonessential region is sufficient.

By the homologous recombination of the above-mentioned homology vector and the parent herpesvirus, a  
15 recombinant herpesvirus can be obtained.

As a concrete method of constructing a recombinant herpesvirus, the following method can be illustrated.

The homology vector is introduced into a herpesvirus-infected cell by electroporation, the calcium phosphate method, a method using lipofectin, a method  
20 using a gene gun and the like. For example, when the parent virus is an avian herpesvirus, the cell to be infected by the herpesvirus is preferably an avian-derived cell such as a chick embryo fibroblast (CEF), a developing chicken egg, a chicken kidney cell etc. The infected cell may be cultured by a commonly used culture method. As a method of introducing the homology vector  
25 into the infected cell, electroporation or a method using lipofectin is preferably adopted for the purpose of obtaining a high introduction efficiency. When the amount of the homology vector (plasmid etc.) to be introduced is in the range of 0.1-1000  $\mu$ g, incidences of forming a recombinant herpesvirus from the homologous regions of the herpesvirus genomic DNA and the homology  
30 vector become high. As a method of selecting only such a recombinant herpesvirus to which such a homology vector has been introduced, the black plaque assay (BPA) method

can be used. In the BPA method, an immunological reaction is carried out using an antibody against the foreign gene, and plaques that expressed the foreign antigen are visualized, that is, an antibody against the foreign gene is used and then an enzyme-labelled secondary antibody is used, and finally a substrate corresponding to the enzyme is used for visualization. By this method, a recombinant herpesvirus that expressed the foreign gene is selected. Furthermore, it has an advantage that, when such a recombinant herpesvirus is constructed, the detection of integration can be easily carried out using a marker gene such as  $\beta$ -galactosidase as the foreign gene. When the  $\beta$ -galactosidase gene is used, a recombinant can be easily isolated by monitoring the expression using Bluo-Gal (Invitrogen) etc. Otherwise, a method such as plaque hybridization may be used to isolate the desired recombinant herpesvirus. By repeating these procedures, recombinant herpesvirus can be purified.

#### Vaccine for infectious laryngotracheitis virus

The vaccine of the present invention for infectious laryngotracheitis virus is a vaccine for avians comprising as an active ingredient the above recombinant herpesvirus of the present invention.

The method of preparing the vaccine is not specifically limited, and may be prepared by, for example, the following method.

The cells infected with the recombinant herpesvirus of the present invention are infected to the cells (hereinafter referred to as the host cell) in which said virus can grow, and after the viruses are grown cells are scraped by a scraper or trypsin, followed by centrifugation to separate the infected cells and the supernatant. For example, when the parent virus is an avian herpesvirus, an avian-derived cell is preferred as the host cell, and chick embryo fibroblasts (CEF), chicken kidney cells etc. can be suitably used. The

infected cells obtained are suspended in a culture medium containing 10% dimethyl sulfoxide (DMSO) and stored frozen in liquid nitrogen. When they are used as the vaccine, a suitable amount of phosphate buffer or physiological saline is used to dissolve the lyophilized product before use.

Stabilizers or other ingredients for storing the above infected cells under liquid nitrogen are any ingredients that allow stable survival of the virus-infected cells and that are not pharmacologically harmful to the recipient.

The method of administering the vaccine comprising as an main ingredient the recombinant herpesvirus thus prepared to avians is not specifically limited. For example, there can be mentioned the same method currently used for herpesvirus vaccines such as a method of injecting subcutaneously to an individual avian and a method of inoculating by puncturating a growing egg.

The amount inoculated and the timing of inoculation may be the same as those used for the current vaccines. For example, by inoculating a dose of  $10^2$ - $10^5$  PFU or  $10^2$ - $10^4$  TCID<sub>50</sub> subcutaneously on the back of an avian on the day of hatching using a 20G or larger needle, the effect as vaccine can be expected. Or the same dose as above may be inoculated by puncturating a developing egg on day 18-19 after development. In addition to the method of administering to avians using a needle as the method of inoculation, an in ovo inoculation instrument such as Inovoject (Embrex) may be used.

The recombinant herpesvirus obtained as above functions not only as a vaccine against ILTV but as a vaccine against the parent herpesvirus as well.

#### Examples

Example 1. Construction of a recombinant plasmid (homology vector) having the full-length ILTV gB gene

A BglIII-cleaved 125 bp fragment of pGHMCSpolyASfi

described in U.S. Pat. Pub. No. 2003-0059799 was inserted into the SfiI fragment of pNZ45/46Sfi described in International Patent Publication WO99/18215 (EP1026246) to construct P45/46HMCSpolyASfi. With pUC18Xlac  
5 described in International Patent Publication WO98/18215 (EP1026246) as the template, PCR was carried out using a primer M13(-21) set forth in SEQ ID NO: 6 and a primer lac3'KpnR set forth in SEQ ID NO: 7 to obtain a 3205 bp fragment.

10 PCR used the Pfu polymerase (Stratagene Corp.) and the DNA Thermal Cycler 480 of Perkin Elmer Inc. and was carried out for 30 cycles under a standard condition (denaturation at 95°C for 1 min, annealing at 60°C or 55°C for 2 minutes, and extension at 72°C for 3 minutes). This  
15 condition was used in all Examples unless otherwise specified.

A 3149 bp fragment obtained by digesting the PCR-amplified 3205 bp fragment with BamHI and KpnI was ligated to a 5573 bp fragment obtained by digesting  
20 P45/46HMCSpolyASfi with BamHI and KpnI to construct pNZ45/46HlacpolyASfi.

On the other hand, with pBK-CMV (Stratagene Corp.) as the template, PCR was carried out using a primer M13(-21) set forth in SEQ ID NO: 6 and a primer pCMV-1 set  
25 forth in SEQ ID NO: 8 to obtain a 953 bp fragment. A 599 bp fragment obtained by digesting this 953 bp fragment with PstI and NheI, a 382 bp fragment obtained by digesting pNZ45/46HlacpolyASfi with PstI and SphI and a 8332 bp fragment obtained by digesting  
30 pNZ45/46HlacpolyASfi with SphI and XbaI were subjected to three-fragment ligation to construct pNZ45/46HCMVlac.

In order to delete the BglII cleavage site in the ILTV gB gene without changing the encoding amino acids, two fragments were obtained with pGTPs/ILgB described in  
35 Japanese Unexamined Patent Publication (Kokai) No. 10-807866 (EP953642) as the template, a 1132 bp fragment obtained by PCR amplification using a primer ILgB-5 set

forth in SEQ ID NO: 9 and a primer ILgB-BglR set forth in  
SEQ ID NO: 10, and a 1564 bp fragment obtained by PCR  
amplification using a primer ILgB-Bgl set forth in SEQ ID  
NO: 11 and a primer ILgB-3+Kpn set forth in SEQ ID NO:  
5 12. With the two fragments as the template, PCR was  
carried out using a primer ILgB-5 set forth in SEQ ID NO:  
9 and a primer ILgB-3+Kpn set forth in SEQ ID NO: 12 to  
obtain a 2648 bp fragment. A 2638 bp fragment obtained  
by digesting this 2648 bp fragment with BamHI and KpnI  
10 and a 3280 bp fragment obtained by digesting pGIPEC  
described in Japanese Unexamined Patent Publication  
(Kokai) No. 2001-000188 with BamHI and KpnI were ligated  
to construct pGIPECILgB.

A 2638 bp fragment obtained by digesting this  
15 pGIPECILgB with BamHI and KpnI and a 4479 bp fragment  
obtained by digesting pGIBacpA described in EP1298139  
with BamHI and KpnI were ligated to construct pGIBAcgBpA.

By inserting a 4464 bp fragment obtained by  
digesting this pGIBAcgBpA with BglI into the SfiI site of  
20 the above pNZ45/46HCMVlac, a homology vector  
p45/46HCMVlacBacgB2nd having CMV-IE as the promoter was  
constructed.

Then, in order to delete the BglI cleavage site in  
the CMV promoter, a primer pCMV-1 set forth in SEQ ID NO:  
25 8 and a primer pPec1R set forth in SEQ ID NO: 13 were  
amplified by PCR with pGIPEC described in Japanese  
Unexamined Patent Publication (Kokai) No. 2001-000188 as  
the template to obtain a 293 bp fragment. With pBK-CMV  
(Stratagene Corp.) as the template, a primer pCMV-01 set  
30 forth in SEQ ID NO: 14 and a primer pCMV-R1 set forth in  
SEQ ID NO: 15 were used in PCR amplification to obtain a  
341 bp fragment. With these two fragments as the  
template, a primer pCMV-1 set forth in SEQ ID NO: 8 and a  
primer pCMV-R1 set forth in SEQ ID NO: 15 were used in  
35 PCR amplification to obtain a 604 bp fragment. A 589 bp  
fragment obtained by digesting this 604 bp fragment with  
PstI and XbaI and a 2765 bp fragment obtained by

digesting pGIPEC described in Japanese Unexamined Patent Publication (Kokai) No. 2001-000188 with PstI and XbaI were ligated to construct pGICMV(-).

5 A 2137 bp fragment obtained by digesting this pGICMV(-) with BamHI and XhoI and a 4054 bp fragment obtained by digesting the above pGIBAcgBpA with BamHI and XhoI were ligated to construct pCMV-ILgB. By inserting a 3338 bp fragment obtained by digesting this pCMV-ILgB with BglI into the SfiI site of pNZ45/46RSVlac-T, a  
10 homology vector p45/46CMVILgBlac, having CMV-IE as the promoter, was constructed.

A 2103 bp fragment obtained by digesting pGIPEC described in Japanese Unexamined Patent Publication (Kokai) No. 2001-000188 with BamHI and XhoI and a 4054 bp  
15 fragment obtained by digesting the above pGIBAcgBpA with BamHI and XhoI were ligated to construct pGIPECILgB2.

By inserting a 3504 bp fragment obtained by digesting this pGIPECILgB2 with BglI into the SfiI site of pNZ45/46RSVlac-T described in Japanese Unexamined  
20 Patent Publication (Kokai) No. 11-192093 (EP1026246), a homology vector p45/46PecILgBlac having the Pec promoter as the promoter was constructed.

Similarly, by inserting a 3504 bp fragment obtained by digesting the above pGIPECILgB2 with BglI into the  
25 SfiI site of pNZ45/46Sfi described in Japanese Unexamined Patent Publication (Kokai) No. 11-192093 (EP1026246), a homology vector p45/46PecILgB having the Pec promoter as the promoter was constructed.

30 Example 2. Construction and purification of a recombinant HVT having a full-length ILTV gB gene

Using the four homology vectors (p45/46HCMVlacBacgB2nd, p45/46CMVILgBlac, p45/46PecILgBlac, p45/46PecILgB) constructed in Example  
1, a recombinant HVT was constructed and purified.

35 Specifically the following procedure was carried out.

First, the DNA of HVT was collected according to the

method of Morgan et al. (Avian Dis., Vol. 34, 345-351, 1990). Thus, about  $1 \times 10^5$  PFU of HVT, the FC126 strain (ATCC VR-584B) or the FC126 strain (the original strain provided by Dr. Witter) was infected to about  $3 \times 10^7$  CEF, and after culturing for 2-3 days, 4 ml of the lysis buffer (0.5% SDS, 10 mM Tris (pH8.0), 100 mM NaCl, 1 mM EDTA, 200 µg/ml proteinase K) was added thereto and incubated at 37°C for 4 hours, followed by phenol extraction and ethanol precipitation to collect HVT-DNA.

10       The homology vectors were cleaved with restriction enzymes at regions containing no homologous sites or foreign genes and linearized.

      About  $3 \times 10^6$  CEF collected with trypsin was suspended into saline G (0.14 M NaCl, 0.2 mM KCl, 1.1 mM disodium hydrogen phosphate, 1.5 mM monobasic potassium phosphate, 0.5 mM magnesium chloride hexahydrate, 0.011% glucose), 10-30 µg of the above HVT-DNA and 10-30µg of a linearized homology vector for recombination were each electroporated using the gene pulser (manufactured by Bio-Rad Inc.) under the condition of  $0.5 \text{ KVcm}^{-1}$  and 0.4 msec. The cell suspension was plated on a culture dish with a diameter of 6 cm, and after adding a culture medium, it was cultured for 5-7 days. After culturing, the virus containing the recombinant HVT was collected from the culture liquid. Then the limiting dilution was carried out to purify recombinant HVT.

      The specific method of purification was as follows:

      For the homology vector p45/46PecILgB containing no lacZ gene, about  $2 \times 10^6$  CEF together with the serially diluted virus solution were plated on 96-well plates. After culturing for 3-5 days, plaques appeared, and a replica was prepared. One of the plates was subjected to screening according to the Black Plaque Assay (BPA) as follows:

35       Antiserum (anti-ILTV-gB antiserum) obtained by immunizing a rabbit with the ILTV-gB protein expressed in

Escherichia coli diluted about 500-fold in a Dulbecco's phosphate buffer (manufactured by Dainippon Seiyaku Co., Ltd.; hereinafter referred to as PBS(-)) containing no magnesium was reacted to plaques at 22-25°C for 2 hours.

5 After washing with 3% non-fat dried milk in PBS(-) three times, it was reacted to plaques at 22-25°C for 2 hours with biotinylated anti-rabbit antibody (sheep, Biosource Inc.). The antibody after the reaction was washed in PBS(-), and then reacted to avidin-biotin-alkaline  
10 phosphatase complex (Vector laboratories). After rinsing off the unreacted avidin-biotin-alkaline phosphatase complex with PBS(-), the color was allowed to change from dark blue to black using BCIP/NBT (manufactured by Roche), a substrate for alkaline phosphatase.

15 In the BPA method, the suspended virus corresponding to positive plaques thus generated is infected again to CEF, and this procedure is repeated for 3-4 times until all plaques become positive with BPA and, usually, the purification and construction of recombinant HVT is  
20 complete.

However, for recombinant HVT that has integrated p45/46PecILgB, recombinant HVT that was purified to 100% could not be obtained.

The construction of recombinant HVT using the  
25 homology vectors p45/46HCMVlacBacgB2nd, p45/46CMVILgBlac and p45/46PecILgBlac containing lacZ was carried out according to the following procedure.

Thus, about  $2 \times 10^6$  CEF together with the serially diluted virus solutions were plated on 96-well multi  
30 plates. After culturing for 3-5 days, plaques appeared, and a replica was prepared. To one of the plates, 100  $\mu$ l/well each of 100  $\mu$ g/ml Blue-gal (manufactured by Gibco), a chromogenic substrate for  $\beta$ -galactosidase, was added, and incubated at 37°C for 4 hours. As plaques that  
35 express lacZ turn blue, cells were collected from those wells in another plate corresponding to wells that



contain blue-colored plaques to make it a virus solution. This virus solution was plated together with about  $2 \times 10^6$  CEF on a flat-bottomed 96-well multi plate for culturing. A step of subculturing from a flat-bottomed 96-well multi  
5 plate for culturing to a flat-bottomed 96-well multi plate for culturing was considered to be one screening. Screening was repeated until all wells had blue plaques, and purification was carried out until all plaques turned blue (usually about 5-10 times of screening) when Blue-  
10 gal was added.

As a result, one clone of recombinant HVT was successfully purified only from the homology vector p45/46PecILgBlac, and this was named FW050.

When p45/46HCMVlacBacgB2nd or p45/46CMVILgBlac was  
15 used, no 100%-purified recombinant HVT was obtained.

Example 3. The structure and vaccine effect of recombinant HVT FW050

FW050, a recombinant HVT obtained in Example 2, was directly sequenced. The result revealed that FW050 has a  
20 deletion of 1542 bp containing the polyA signal, and instead three bases (GCG) of unknown origin have been sandwiched (SEQ ID NO: 22). As a result, it was predicted to express a chimera protein consisting of 469  
25 amino acids that comprises 429 amino acids at the amino terminal end of the ORF of the original ILTV gB gene and 40 amino acids that are now encoded by the polyA signal portion until the stop codon appears (see Fig. 1 and SEQ ID NO: 1).

This FW050 was subjected to an animal experiment for  
30 examining the vaccine effect and the result shown in Table 1 was obtained.

The animal experiment was carried out according to the provisions (9CFR) in the Animal and Plant Health Inspection Service (APHIS) by the Department of  
35 Agriculture of the USA.

Challenge of an highly toxic ILTV was carried out using a method described in 9CFR Ch.1 113.328. Thus, a

recombinant HVT was inoculated to over 10 SPF chickens per group (Line M: Nippon Institute for Biological Science) (In Experiment 1 alone, the number of inoculated chickens for FW050 was 9 due to the accidental death of one animal after inoculation). The negative control group (control) received no inoculation.

When the test SPF chickens hatched, recombinant HVT was subcutaneously inoculated to the back of chickens at  $1 \times 10^4$  PFU using a 26G needle. To 4 week-old animals, a highly virulent ILTV (the NS-175 strain,  $1 \times 10^{3.0}$  EID<sub>50</sub>/0.1 ml) was challenge-inoculated to the infraorbital groove. For ten days after inoculation, clinical symptoms were daily observed and the presence or absence of infection prevention was determined.

The result is shown in Table 1.

As can be seen from Table 1, a protective effect against the highly virulent ILTV strain was recognized for FW050.

Table 1

|           | Recombinant HVT | Homology vector  | Protection rate % (No. of chickens protected/total No. of chickens) |              |
|-----------|-----------------|------------------|---|--------------|
|           |                 |                  | Experiment 1  | Experiment 2 |
| Control   |                 |                  | 0 (0/10)  | 0 (0/16)     |
| Challenge | FW050           | p45/46PecILgBlac | 78 (7/9)  | 31 (4/13)    |

Example 4. Construction of a recombinant having a fragment in which the gB gene of ILTV was truncated

A homology vector having a mutant gB gene product in which part of ORF of the ILTV gB gene was deleted was constructed as follows:

(1) A homology vector that contains DNA encoding 623-amino acid gB-a from the amino terminal that is the ORF of the ILTV gB gene and that is unlikely to contain a dimer-forming region: p45/46PecILgBa and p45/46PecILgBalac

Using pGIPECILgB2 constructed in Example 1 as the template, a 1205 bp fragment was amplified by the above-

mentioned standard PCR using a primer P-BglIII set forth in SEQ ID NO: 16 and a primer A-R set forth in SEQ ID NO: 17. A 1198 bp fragment obtained by digesting this with BglIII and KpnI and a 7057 bp fragment obtained by digesting p45/46PecILgB constructed in Example 1 with BglIII and KpnI were ligated to construct a homology vector p45/46PecILgBa.

A 6373 bp fragment obtained by digesting p45/46PecILgBa with BglIII and XhoI and a 5923 bp fragment obtained by digesting p45/46PecILgBlac constructed in Example 1 with BglIII and XhoI were ligated to construct a homology vector p45/46PecILgBalac.

(2) A homology vector that contains DNA encoding 691-amino acid gB-b in which the carboxy terminal has been deleted to immediately before the transmembrane domain: p45/46PecILgBb and p45/46PecILgBblac

Using pGIPecILgB2 constructed in Example 1 as the template, a 1409 bp fragment was amplified by the above-mentioned standard PCR using a primer P-BglIII set forth in SEQ ID NO: 16 and a primer B-R set forth in SEQ ID NO: 18. A 1402 bp fragment obtained by digesting this with BglIII and KpnI and a 7057 bp fragment obtained by digesting p45/46PecILgB constructed in Example 1 with BglIII and KpnI were ligated to construct a homology vector p45/46PecILgBb.

A 6577 bp fragment obtained by digesting p45/46PecILgBb with BglIII and XhoI and a 5923 bp fragment obtained by digesting p45/46PecILgBlac constructed in Example 1 with BglIII and XhoI were ligated to construct a homology vector p45/46PecILgBblac.

(3) A homology vector that contains DNA encoding 803-amino acid gB-c in which the transmembrane domain alone has been deleted: p45/46PecILgBc and p45/46PecILgBclac

Using pGIPecILgB2 constructed in Example 1 as the template, a 1409 bp fragment was amplified by the above-mentioned standard PCR using a primer P-BglIII set forth

in SEQ ID NO: 16 and a primer C-R set forth in SEQ ID NO: 19. Similarly, using pGI<sub>PecILgB2</sub> as the template, a 357 bp fragment was amplified by the standard PCR using a primer C-F set forth in SEQ ID NO: 20 and a primer CDE-R set forth in SEQ ID NO: 21. Using two fragments of the 1409 bp fragment and the 357 bp fragment as the template, a 1745 bp fragment was amplified by the standard PCR using a primer P-BglIII set forth in SEQ ID NO: 16 and a primer CDE-R set forth in SEQ ID NO: 21. A 1738 bp fragment obtained by digesting this with BglIII and KpnI and a 7057 bp fragment obtained by digesting p45/46<sub>PecILgB</sub> constructed in Example 1 with BglIII and KpnI were ligated to construct a homology vector p45/46<sub>PecILgBc</sub>.

A 6913 bp fragment obtained by digesting p45/46<sub>PecILgBc</sub> with BglIII and XhoI and a 5923 bp fragment obtained by digesting p45/46<sub>PecILgBlac</sub> constructed in Example 1 with BglIII and XhoI were ligated to construct a homology vector p45/46<sub>PecILgBclac</sub>.

The schematic diagrams of these homology vectors are shown in Fig. 2.

Using these six homology vectors, thus constructed, purification of recombinant HVTs were attempted in a manner similar to that described in Example 2, but recombinants having the same structure as the homology vectors could not be obtained.

Thus, though a plurality of homology vectors were constructed that were truncated from the carboxy terminal of the gB protein subsequent to the promoter to purify and construct recombinant HVT, recombinants having the same length of the ILTV gB gene as the homology vectors could not be obtained. This result demonstrated that when the gB gene integrated together with the promoter was to be expressed in recombinant HVT, those having a long ORF cannot be purified. Also, even if recombinant HVT could be purified, it was thought, the ORF of the gB gene in the recombinant HVT obtained becomes curtailed

due to selection pressure. Thus, it was predicted that when the gB gene is integrated together with the promoter, the length of gB gene ORF of ILTV that can be present as a stable recombinant HVT is about the size of those encoding 623 amino acids or shorter.

Example 5. Modification from the deleted clone FW050 was reverted and the vaccine effect of the recombinant HVT

From the recombinant HVT FW050 for which the vaccine effect was confirmed in Example 3 and which is expected to express a fusion protein of the 429 amino acids from the amino terminal end of the gB protein of ILTV and the 40 amino acids derived from polyA, a homology vector p45/46CMVILgBf was constructed in order to express a protein in which 40 amino acids derived from polyA of the protein to be expressed have been deleted.

Specifically, a 303 bp fragment obtained by digesting pGIBacpA described in Japanese Unexamined Patent Publication (Kokai) No. 2004-000111 with EcoRI and KpnI and a 5854 bp fragment obtained by digesting pGIPecILgB2 constructed in Example 1 with EcoRI and KpnI were ligated to construct pGIPecILgB3. A 2245 bp fragment obtained by digesting this pGIPecILgB3 with BglII and SfiI and a 6759 bp fragment obtained by digesting p45/46PecILgB with BglII and SfiI were ligated to construct p45/46PecILgB2.

Then, using pGIPecILgB2 as the template, P-BglII set forth in SEQ ID NO: 16 and F-R set forth in SEQ ID NO: 23 were subjected to PCR to obtain a 623 bp fragment. A 616 bp fragment obtained by digesting this with BglII and KpnI and a 7057 bp fragment obtained by digesting p45/46PecILgB2 with BglII and KpnI were ligated to construct p45/46PecILgBf.

A 7102 bp fragment obtained by complete digestion of this p45/46PecILgBf with XbaI followed by partial digestion with PstI and a 589 bp fragment obtained by digesting pGICMV(-) constructed in Example 1 with PstI and XbaI were ligated to complete the construction of a

homology vector p45/46CMVILgBf.

Using this homology vector p45/46CMVILgBf, FW063, a recombinant HVT, was constructed and purified. It was confirmed that the inserted gene portion of the recombinant HVT FW063 was the same as the base sequence of the homology vector and had not been mutated.

Furthermore, by reversely cloning FW050, a homology vector p45/46PecILgBdellac was constructed in the following procedure.

The template used in PCR for this cloning is a viral DNA collected from a chicken that was inoculated with FW050, and the primer was two primers of P-BglII set forth in SEQ ID NO: 16 and 45/46F(K) set forth in SEQ ID NO: 24, and the homology vector p45/46PecILgBdellac was constructed by ligating a 707 bp fragment obtained by digesting, with BglII and SfiI, a 944 bp fragment obtained by amplification with the two primers and a 10809 bp fragment obtained by digesting p45/46PecILgBelac with BglII and SfiI. Using this homology vector p45/46PecILgBdellac, recombinant HVT FW069 was successfully purified and constructed.

In the same manner as above, without changing the polyA portion at all, a homology vector p45/46PecILgBdellac+STP in which termination was inserted into a site where the ORF of 429 amino acids at the amino terminal end of the gB protein terminated was constructed in the following procedure.

First, using, as the template, a 637 bp fragment amplified using a primer P-BglII set forth in SEQ ID NO: 16 and a primer gBdelSTP-R set forth in SEQ ID NO: 25 and a 673 bp fragment amplified using a primer gBdelSTP-F set forth in SEQ ID NO: 26 and a primer 45/46F(B) set forth in SEQ ID NO: 27, PCR was carried out with the primer P-BglII and the primer 45/46F(B) to obtain a 1263 bp fragment. Then a 707 bp fragment obtained by digesting this with BglII and SfiI and a 10809 bp fragment obtained by digesting p45/46PecILgBlac with BglII and SfiI were

ligated to construct a homology vector  
p45/46PecILgBdellac+STP.

Using this homology vector p45/46PecILgBdellac+STP,  
a recombinant HVT FW070 was successfully purified and  
constructed.

Thus, FW063, FW069 and FW070, in which the ORF of  
the gB gene is 429 amino acids as in FW050 described in  
Example 3, could easily be purified. This demonstrated  
that when DNA derived from the gB gene encoding the 429  
amino acids at the amino terminal end of the amino acid  
sequence set forth in SEQ ID NO: 2 is to be expressed in  
a recombinant HVT that has integrated it together with  
the promoter, purification can be carried out.

Using the recombinant HVT thus purified, FW063,  
FW069 and FW050 purified in Example 3, a challenge test  
was carried out as in Example 3 to investigate the  
vaccine effect. The result is shown in Table 2.

Table 2

|           | Recombinant<br>HVT | Homology vector     | Protection rate % (No.<br>of chickens protected /<br>total No. of chickens) |
|-----------|--------------------|---------------------|---|
| Control   |                    |                     | 0 (0/7)   |
| Challenge | FW050              | p45/46PecILgBlac    | 54 (7/13)   |
| Challenge | FW063              | p45/46CMVILgBf      | 33 (5/15)   |
| Challenge | FW069              | p45/46PecILgBdellac | 56 (9/16)   |

From the result, it can be seen that the vaccine  
effect can also be recognized for FW063 having no  
additive sequence, and FW069 and FW050 that have the  
additive sequence exhibit a better vaccine effect.